

FIG 1

ADAMTS-E nucleotide sequence [SEQ ID NO: 1]

CACGCGTCCGACGGCGCGGAGGCCCGGGCGCGGCGCAGGAGCCCGGTGAT
5 GCTGCGAAGGCTGTGAACAGGGGAGGCGGCACTGTGGGGGCTGCCGGCAGCCGGGG
CTGGGGAGAGACATGTGGACACGTGGCCTCTATGGCTCCCGCCTGCCAGATCCTCCGC
TGGGCCCTCGCCCTGGGGCTGGGCCTCATGTTGAGGTCACGCACGCCTTCCGGTCTC
AAGATGAGTTCCTGTCCAGTCTGGAGAGCTATGAGATCGCCTTCCCCACCCGCGTGGAC
CACAACGGGGCACTGCTGGCCTTCTCGCCACCTCCTCCCCGGAGGCAGCGCCGCGGC
10 ACGGGGGCCACAGCCGAGTCCCGCCTCTTCTACAAAGTGGCCTCGCCAGCACCCACT
TCCTGCTGAACCTGACCCGCAGCTCCCGTCTACTGGCAGGGCACGTCTCCGTGGAGTA
CTGGACACGGGAGGGCCTGGCCTGGCAGAGGGCGGCCCGGCCCACTGCCTCTACGC
TGGTCACCTGCAGGGCCAGGCCAGCACCTCCCATGTGGCCATCAGCACCTGTGGAGGC
CTGCACGGCCTGATCGTGGCAGACGAGGAAGAGTACCTGATTGAGCCCTGCACGGTG
15 GGCCCAAGGGTTCTCGGAGCCCGGAGGAAAGTGGACCACATGTGGTGTACAAGCGTTC
CTCTCTGCGTCACCCCCACCTGGACACAGCCTGTGGAGTGAGAGATGAGAAACCGTGG
AAAGGGCGGCCATGGTGGCTGCGGACCTTGAAGCCACCGCCTGCCAGGCCCTGGGG
AATGAAACAGAGCGTGGCCAGCCAGGCCTGAAGCGATCGGTGAGCCGAGAGCGCTACG
TGGAGACCCTGGTGGTGGCTGACAAGATGATGGTGGCCTATCACGGGCGCCGGGATGT
20 GGAGCAGTATGTCCTGGCCATCATGAACATTGTTGCCAACTTTTCCAGGACTCGAGTCT
GGGAAGCACCGTTAACATCCTCGTAACTCGCCTCATCCTGCTCACGGAGGACCAGCCCA
CTCTGGAGATCACCCACCATGCCGGGAAGTCCCTGGACAGCTTCTGTAAGTGGCAGAAA
TCCATCGTGAACCACAGCGGCCATGGCAATGCCATTCCAGAGAACGGTGTGGCTAACCA
TGACACAGCAGTGCTCATCACACGCTATGACATCTGCATCTACAAGAACAACCCTGCG
25 GCACACTAGGCCTGGCCCCGGTGGGCGGAATGTGTGAGCGCGAGAGAAGCTGCAGCG
TCAATGAGGACATTGGCCTGGCCACAGCGTTCACCATTGCCACGAGATCGGGCACACA
TTCGGCATGAACCATGACGGCGTGGGAAACAGCTGTGGGGCCCGTGGTCAGGACCCAG
CCAAGCTCATGGCTGCCACATTACCATGAAGACCAACCCATTCTGTGTGGTCATCCTGC
AGCCGTGACTACATCACCAGCTTTCTAGACTCGGGCCTGGGGCTCTGCCTGAACAACCG
30 GCCCCCCAGACAGGACTTTGTGTACCCGACAGTGGCACCGGGCCAAGCCTACGATGCA
GATGAGCAATGCCGCTTTCAGCATGGAGTCAAATCGCGTCAGTGTAATACGGGGAGGT
CTGCAGCGAGCTGTGGTGTCTGAGCAAGAGCAACCGGTGCATCACCAACAGCATCCCG
GCCGCCGAGGGCACGCTGTGCCAGACGCACACCATCGACAAGGGGTGGTGTACAAAC
GGGTCTGTGTCCCCTTTGGGTGCGGCCAGAGGGTGTGGACGGAGCCTGGGGGCCGT
35 GGACTCCATGGGGCGACTGCAGCCGACCTGTGGCGGCGGCGTGTCTCTTAGCC

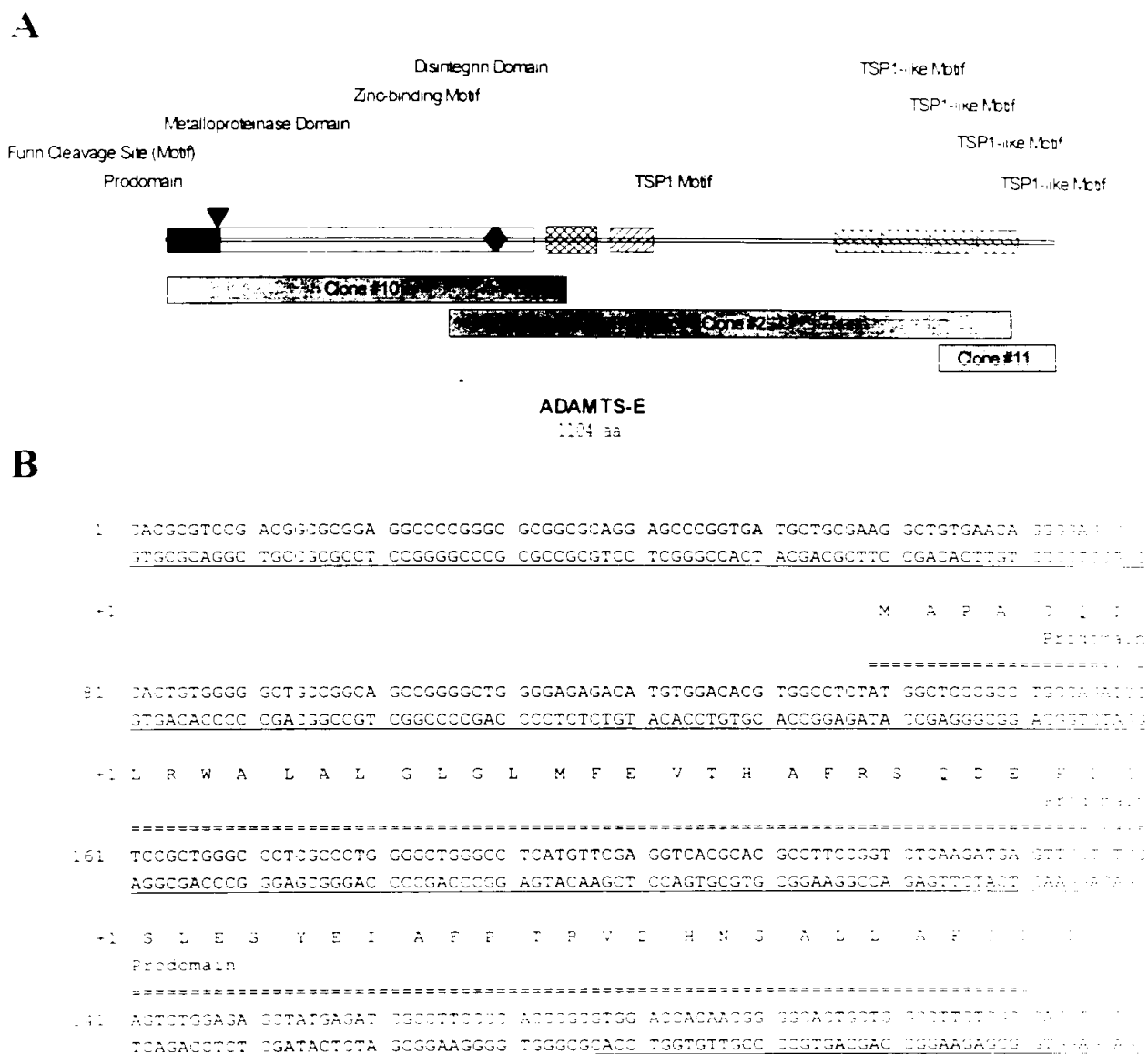
GTGCAGTGTCTGAATTTGACAGCATCCCTTTCCGTGGGAAATTCTACAAGTGGAAAACG
TACCGGGGAGGGGGCGTGAAGGCCTGCTCGCTCACGTGCCTAGCGGAAGGCTTCAACT
TCTACACGGAGAGGGCGGCAGCCGTGGTGGACGGGACACCCTGCCGTCCAGACACGG
TGGACATTTGCGTCAGTGGCGAATGCAAGCACGTGGGCTGCGACCGAGTCCTGGGCTC
5 CGACCTGCGGGAGGACAAGTGCCGAGTGTGTGGCGGTGACGGCAGTGCCTGCGAGAC
CATCGAGGGCGTCTTCAGCCAGCCTCACCTGGGGCCGGGTACGAGGATGTCGTCTGG
ATTCCCAAAGGCTCCGTCCACATCTTCATCCAGGATCTGAACCTCTCTCTCAGTCACTTG
GCCCTGAAGGGAGACCAGGAGTCCCTGCTGCTGGAGGGGCTGCCCCGGGACCCCCCAG
CCCCACCGTCTGCCTCTAGCTGGGACCACCTTTCAACTGCGACAGGGGGCCAGACCAGG
10 TCCAGAGCCTCGAAGCCCTGGGACCGATTAAATGCATCTCTCATCGTCATGGTGCTGGCC
CGGACCGAGCTGCCTGCCCTCCGCTACCGCTTCAATGCCCCCATCGCCCGTGA CTGCG
TGCCCCCTACTCCTGGCACTATGCGCCCTGGACCAAGTGCTCGGCCCAAGTGTGCAGG
CGGTAGCCAGGTGCAGGCGGTGGAGTGCCGCAACCAGCTGGACAGCTCCGCGGTGCG
CCCCCACTACTCCAGTGCCCAACAGCAAGCTGCCCCAAAAGGCAGCGCGCCTGCAACACG
15 GAGCCTTGCCCTCCAGACTGGGTTGTAGGGAAGTGGTCGCTCTGCAGCCGCAGCTGCG
ATGCAGGCGTGCGCAGCCGCTCGGTCTGTGTGCCAGCGCCGCGTCTCTGCCGCGGAGG
AGAAGGCGCTGGACGACAGCGCATGCCCCGAGCCGCGCCACCTGTACTGGAGGCCT
GCCACGGCCCCACTTGCCCTCCGGAGTGGGCGGCCCTCGACTGGTCTGAGTGCACCC
CCAGCTGCGGGGCCGGGCCTCCGCCACCGCGTGGTCCTTTGCAAGAGCGCAGACCACC
20 GCGCCACGCTGCCCCCGGCGCACTGCTCACCCGCGCCAAAGCCACCGGCCACCATGC
GCTGCAACTTGCGCCGCTGCCCCCGGCCGCTGGGTGGCTGGCGAGTGGGGTGAGT
GCTCTGCACAGTGCGGCGTCGGGCAGCGGCAGCGCTCGGTGCGCTGCACCAGCCACA
CGGGCCAGGCGTCGCACGAGTGCACGGAGGCCCTGCGGCCGCGGACTACCACGCAGC
AGTGTGAGGCCAAGTGCAGACAGCCCAACCCCCGGGGACGGCCCTGAAGAGTGCAAGG
25 ATGTGAACAAGTTCGCCTACTGCCCCCTGGTGCTCAAATTTAGTTCTGCAGCCGAGCC
TACTTCCGCCAGATGTGCTGCAAAACCTGCCAGGGGCCACTAGGGGGCGCGCGGCACCC
GGAGCCACAGCTGGCGGGGTCTCCGCCGCCAGCCCTGCAGCGGGCCGGCCAGAGGG
GGCCCCGGGGGGGGGCGGGAAGTGGGAGGGGAAGGGTGAGACGGAGCCGGAAGTTATTT
ATTGGGAACCCCTGCAGGGCCCTGGCTGGGAGGATCCACCCCAACCTCTGCCCTGCCC
30 GCCCCAGGGGCACCCCGACATCCAGGCCACCCCTCATGGTGCTACAGACCCTGCCCT
GGGGCCACACACTCCTGCCAGGAAGCCCTACATCAATAAAGTTCTGTCTTGTGTAGAT
TTCTAAAAAAAAAAAAAAAA

FIG 2

ADAMTS-E amino acid sequence [SEQ ID NO: 2]

MAPACQILRWALALGLGLMFEVTHAFRSQDEFLSSLESYEIAFPTRVDHNGALLAFS
PPPPRRQRRGTGATAESRLFYKVASPSTHFLNLTRSSRLLAGHVSVEYWTREGLA
5 WQRAARPHCLYAGHLQGQASTSHVAISTCGGLHGLLIVADEEEYLIPLHGGPKGSR
SPEESGPHVVYKRSSLRHPHLDTACGVRDEKPKWGRPWWLRTLKPPPARPLGNE
TERGQPGLKRSVSRERYVETLVVADKMMVAYHGRRDVEQYVLAIMNIVAKLFQDSS
LGSTVNILVTRLILLTEDQPTLEITHHAGKSLDSFCKWQKSIVNHSGHGNAIPENGVA
NHDTAVLITRYDICIYKNKPCGTLGLAPVGGMCERERSCSVNEDIGLATAFTIAHEIG
10 HTFGMNHGDBGVNSCGARGQDPAKLMAAHITMKTNPFWSSCSRDIYTSFLDSGLG
LCLNNRPPRQDFVYPTVAPGQAYDADEQCRFQHGKSRQCKYGEVCSSELWCLSK
SNRCITNSIPAAEGTLCQHTIDKGWCYKRVCPFGSRPEGVDGAWGPWTPWGDG
SRTCAGGVSSSRHCDSRPTIGGKYCLGERRRHRSCNTDDCPPGSQDFREVQC
SCFDSIPFRCKFYKWKTYRGGGVKACSLTCLAEGFNFYTERAAAVVDGTPCRPDTV
15 DICVSGECKHVGCDRVLGSDLREDKCRVCGGDGSACETIEGVFSPASPGAGYEDV
VWIPKGSVHIFIQDLNLSLSHLALKGDQESLLEGLPGTPQPHRLPLAGTTFFQLRQGP
DQVQSLEALGPINASLIVMVLARTELPALRYRFNAPIARDSLPPYSWHYAPWTKCSA
QCAGGSQVQAVECRNQLDSSAVAPHYCSAHSKLPKRQRACNTEPCPPDWVVG
WSLCSRSCDAGVRSRSVVCQRRVSAAEEKALDDSACPQPRPPVLEACHGPTCPPE
20 WAALDWSECTPSCGPGLRHRVVLCKSADHRATLPPAHCSPAAPKPPATMRCNLRR
PPARWWAGEWGECSAQCGVGQRQSVRCTSHTGQASHECTEALRPPTTTTQQCE
AKCDSPTPGDGPEECKDVNKVAYCPLVLKFQFCSRAYFRQMCKKTCQGH

Figure 3. Domain structure of ADAMTS-E and translated nucleic acid sequence. A) Diagram of ADAMTS-E showing the following domains and signature motifs (with amino acid numbers in parentheses): prodomain (1-66), furin cleavage site (63-66), metalloproteinase domain (67-453), zinc-binding motif (392-420), disintegrin domain (469-531), TSP1 motif (548-601), and four TSP1-like motifs (829-884, 888-944, 948-1002, and 1007-1058). Overlapping clones covering the indicated sequence segments are depicted at the bottom of the diagram. **B)** ADAMTS-E nucleotide sequence with translated amino acid sequence above.



[illegible]

... L I F A A K V I C K L W J L D F C N R I I I
Disintegrin domain

=====
174 TAA TATAA ATATATGAA GTTGTGATG AGTTGTGTTG TGTGATGAA AGTATATG ATATTA ATATTA
TGTGATGAT TATATATG TATATATG TATATATG TATATATG TATATATG TATATATG TATATATG

... A A E I T L I L T R I I C F D W I I F R I
Disintegrin domain

=====
177 ATGTGTGAGG GTAGGTGTG GTAGATGAGG AGCATGAGG AGGGGTGTG GTAGAAATGT GTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... R R E G V D G A W S P W T P W G D C S R T I R

=====
179 GGGGGGAGG GTGTGTGTGT GTAGATGAGG GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... S S S S R H C D S P R P T I G G K Y D L S E R R V I R
TSPI Motif

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181 GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... S C N T D D C P P G S Q D F R E V Q C S E F D E I I R
TSPI Motif

=====
183 TGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... R G K F Y K W K T Y R G G G V K A C S L T C L A E V
185 GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... F N F Y T E R A A A V V D G T P C R P D T V D I T V R
187 TGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... G E C K H V G C D R V L G S D L R E D K C P V T S V R
189 GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... G S A C E T I E G V F S P A S P G A G Y E I
191 GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... P K G S V H I F I Q D L N L S L S H L A L K S D L H R
193 GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT
GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT GTGTGTGTGT

... L L L E G L P G T P Q P H R L P L A G T T F L L R I

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100
TSP1-like Motif

1001 TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA
TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100
TSP1-like Motif

1001 TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA
TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA TATTAATTA

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100
1001 AAGTCTGGCT ACTGCCCCCT GGTGCTCAAA TTTCAGTTCT SCAGCCGAGC CTACTTCCCG CAGATCTAT TAAAAA
TTCTAGCGGA TGACGGGGGA CCACGAGTTT AAAGTCAAGA CTTGGGCTCG CATGAAGGGG CTCTAATTA TTTT

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

1001 CCAGGGGAGC TAGGGGGGCG GCGGCACCCG GAGCCACAGC TGGCGGGGTC TCGGGGGCCA GCGGTGTAAT TATTAATTA
GTTGGGGGTC ATCCCCCGCG CGCCGTGGGC CTCGGTGTCT ACCGCCCCAG AGCGGGCGGT CCGGACCTCT TATTAATTA

1001 GAGGGGGGCG GGGGGGGGCG GGAAGTGGG AGGGAAGGGT GAGACGGAGC CGGAAGTTAT TTATTGGGAA TATTAATTA
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1001 GCGGTGGGTC GGAGGATCCA CCGCAACCTC TCGCCTGCCC GCGCCAGGGG CACCCCGACA TCGAGGGCAC TCGGTATTA
CGGACCGGAC CCGCTAGGT GGGGTTGGAG ACGGGACGGG CGGGGTCCCC CTGGGGCTGT AGGTGGGTC TCGGTATTA

1001 TGCTACAGAC CCTGCCCTGG GCGCCACACA CTCCTGCCAG GAAGCCCTAC ATCAATAAAG TTCTGTCTCT TATTAATTA
ACGATGTCTG GGACGGGACC CCGGGTGTCT GAGGACGGTC CTTCGGGATG TAGTTATTTT AAGATAGAA TATTAATTA

1001 TAAAAAAAAA AAAAAA
ATTTTTTTTT TTTTTT

Metalloproteinase Domain Alignment of ADAMTS-E v. ADAMTS Family

4	45-14-099	6	6-14-099	14	14-14-099	24	24-14-099	34	34-14-099	44	44-14-099	54	54-14-099	64	64-14-099	74	74-14-099	84	84-14-099	94	94-14-099	104	104-14-099	114	114-14-099	124	124-14-099	134	134-14-099	144	144-14-099	154	154-14-099	164	164-14-099	174	174-14-099	184	184-14-099	194	194-14-099	204	204-14-099	214	214-14-099	224	224-14-099	234	234-14-099	244	244-14-099	254	254-14-099	264	264-14-099	274	274-14-099	284	284-14-099	294	294-14-099	304	304-14-099	314	314-14-099	324	324-14-099	334	334-14-099	344	344-14-099	354	354-14-099	364	364-14-099	374	374-14-099	384	384-14-099	394	394-14-099	404	404-14-099	414	414-14-099	424	424-14-099	434	434-14-099	444	444-14-099	454	454-14-099	464	464-14-099	474	474-14-099	484	484-14-099	494	494-14-099	504	504-14-099	514	514-14-099	524	524-14-099	534	534-14-099	544	544-14-099	554	554-14-099	564	564-14-099	574	574-14-099	584	584-14-099	594	594-14-099	604	604-14-099	614	614-14-099	624	624-14-099	634	634-14-099	644	644-14-099	654	654-14-099	664	664-14-099	674	674-14-099	684	684-14-099	694	694-14-099	704	704-14-099	714	714-14-099	724	724-14-099	734	734-14-099	744	744-14-099	754	754-14-099	764	764-14-099	774	774-14-099	784	784-14-099	794	794-14-099	804	804-14-099	814	814-14-099	824	824-14-099	834	834-14-099	844	844-14-099	854	854-14-099	864	864-14-099	874	874-14-099	884	884-14-099	894	894-14-099	904	904-14-099	914	914-14-099	924	924-14-099	934	934-14-099	944	944-14-099	954	954-14-099	964	964-14-099	974	974-14-099	984	984-14-099	994	994-14-099	1004	1004-14-099	1014	1014-14-099	1024	1024-14-099	1034	1034-14-099	1044	1044-14-099	1054	1054-14-099	1064	1064-14-099	1074	1074-14-099	1084	1084-14-099	1094	1094-14-099	1104	1104-14-099	1114	1114-14-099	1124	1124-14-099	1134	1134-14-099	1144	1144-14-099	1154	1154-14-099	1164	1164-14-099	1174	1174-14-099	1184	1184-14-099	1194	1194-14-099	1204	1204-14-099	1214	1214-14-099	1224	1224-14-099	1234	1234-14-099	1244	1244-14-099	1254	1254-14-099	1264	1264-14-099	1274	1274-14-099	1284	1284-14-099	1294	1294-14-099	1304	1304-14-099	1314	1314-14-099	1324	1324-14-099	1334	1334-14-099	1344	1344-14-099	1354	1354-14-099	1364	1364-14-099	1374	1374-14-099	1384	1384-14-099	1394	1394-14-099	1404	1404-14-099	1414	1414-14-099	1424	1424-14-099	1434	1434-14-099	1444	1444-14-099	1454	1454-14-099	1464	1464-14-099	1474	1474-14-099	1484	1484-14-099	1494	1494-14-099	1504	1504-14-099	1514	1514-14-099	1524	1524-14-099	1534	1534-14-099	1544	1544-14-099	1554	1554-14-099	1564	1564-14-099	1574	1574-14-099	1584	1584-14-099	1594	1594-14-099	1604	1604-14-099	1614	1614-14-099	1624	1624-14-099	1634	1634-14-099	1644	1644-14-099	1654	1654-14-099	1664	1664-14-099	1674	1674-14-099	1684	1684-14-099	1694	1694-14-099	1704	1704-14-099	1714	1714-14-099	1724	1724-14-099	1734	1734-14-099	1744	1744-14-099	1754	1754-14-099	1764	1764-14-099	1774	1774-14-099	1784	1784-14-099	1794	1794-14-099	1804	1804-14-099	1814	1814-14-099	1824	1824-14-099	1834	1834-14-099	1844	1844-14-099	1854	1854-14-099	1864	1864-14-099	1874	1874-14-099	1884	1884-14-099	1894	1894-14-099	1904	1904-14-099	1914	1914-14-099	1924	1924-14-099	1934	1934-14-099	1944	1944-14-099	1954	1954-14-099	1964	1964-14-099	1974	1974-14-099	1984	1984-14-099	1994	1994-14-099	2004	2004-14-099	2014	2014-14-099	2024	2024-14-099	2034	2034-14-099	2044	2044-14-099	2054	2054-14-099	2064	2064-14-099	2074	2074-14-099	2084	2084-14-099	2094	2094-14-099	2104	2104-14-099	2114	2114-14-099	2124	2124-14-099	2134	2134-14-099	2144	2144-14-099	2154	2154-14-099	2164	2164-14-099	2174	2174-14-099	2184	2184-14-099	2194	2194-14-099	2204	2204-14-099	2214	2214-14-099	2224	2224-14-099	2234	2234-14-099	2244	2244-14-099	2254	2254-14-099	2264	2264-14-099	2274	2274-14-099	2284	2284-14-099	2294	2294-14-099	2304	2304-14-099	2314	2314-14-099	2324	2324-14-099	2334	2334-14-099	2344	2344-14-099	2354	2354-14-099	2364	2364-14-099	2374	2374-14-099	2384	2384-14-099	2394	2394-14-099	2404	2404-14-099	2414	2414-14-099	2424	2424-14-099	2434	2434-14-099	2444	2444-14-099	2454	2454-14-099	2464	2464-14-099	2474	2474-14-099	2484	2484-14-099	2494	2494-14-099	2504	2504-14-099	2514	2514-14-099	2524	2524-14-099	2534	2534-14-099	2544	2544-14-099	2554	2554-14-099	2564	2564-14-099	2574	2574-14-099	2584	2584-14-099	2594	2594-14-099	2604	2604-14-099	2614	2614-14-099	2624	2624-14-099	2634	2634-14-099	2644	2644-14-099	2654	2654-14-099	2664	2664-14-099	2674	2674-14-099	2684	2684-14-099	2694	2694-14-099	2704	2704-14-099	2714	2714-14-099	2724	2724-14-099	2734	2734-14-099	2744	2744-14-099	2754	2754-14-099	2764	2764-14-099	2774	2774-14-099	2784	2784-14-099	2794	2794-14-099	2804	2804-14-099	2814	2814-14-099	2824	2824-14-099	2834	2834-14-099	2844	2844-14-099	2854	2854-14-099	2864	2864-14-099	2874	2874-14-099	2884	2884-14-099	2894	2894-14-099	2904	2904-14-099	2914	2914-14-099	2924	2924-14-099	2934	2934-14-099	2944	2944-14-099	2954	2954-14-099	2964	2964-14-099	2974	2974-14-099	2984	2984-14-099	2994	2994-14-099	3004	3004-14-099	3014	3014-14-099	3024	3024-14-099	3034	3034-14-099	3044	3044-14-099	3054	3054-14-099	3064	3064-14-099	3074	3074-14-099	3084	3084-14-099	3094	3094-14-099	3104	3104-14-099	3114	3114-14-099	3124	3124-14-099	3134	3134-14-099	3144	3144-14-099	3154	3154-14-099	3164	3164-14-099	3174	3174-14-099	3184	3184-14-099	3194	3194-14-099	3204	3204-14-099	3214	3214-14-099	3224	3224-14-099	3234	3234-14-099	3244	3244-14-099	3254	3254-14-099	3264	3264-14-099	3274	3274-14-099	3284	3284-14-099	3294	3294-14-099	3304	3304-14-099	3314	3314-14-099	3324	3324-14-099	3334	3334-14-099	3344	3344-14-099	3354	3354-14-099	3364	3364-14-099	3374	3374-14-099	3384	3384-14-099	3394	3394-14-099	3404	3404-14-099	3414	3414-14-099	3424	3424-14-099	3434	3434-14-099	3444	3444-14-099	3454	3454-14-099	3464	3464-14-099	3474	3474-14-099	3484	3484-14-099	3494	3494-14-099	3504	3504-14-099	3514	3514-14-099	3524	3524-14-099	3534	3534-14-099	3544	3544-14-099	3554	3554-14-099	3564	3564-14-099	3574	3574-14-099	3584	3584-14-099	3594	3594-14-099	3604	3604-14-099	3614	3614-14-099	3624	3624-14-099	3634	3634-14-099	3644	3644-14-099	3654	3654-14-099	3664	3664-14-099	3674	3674-14-099	3684	3684-14-099	3694	3694-14-099	3704	3704-14-099	3714	3714-14-099	3724	3724-14-099	3734	3734-14-099	3744	3744-14-099	3754	3754-14-099	3764	3764-14-099	3774	3774-14-099	3784	3784-14-099	3794	3794-14-099	3804	3804-14-099	3814	3814-14-099	3824	3824-14-099	3834	3834-14-099	3844	3844-14-099	3854	3854-14-099	3864	3864-14-099	3874	3874-14-099	3884	3884-14-099	3894	3894-14-099	3904	3904-14-099	3914	3914-14-099	3924	3924-14-099	3934	3934-14-099	3944	3944-14-099	3954	3954-14-099	3964	3964-14-099	3974	3974-14-099	3984	3984-14-099	3994	3994-14-099	4004	4004-14-099	4014	4014-14-099	4024	4024-14-099	4034	4034-14-099	4044	4044-14-099	4054	4054-14-099	4064	4064-14-099	4074	4074-14-099	4084	4084-14-099	4094	4094-14-099	4104	4104-14-099	4114	4114-14-099	4124	4124-14-099	4134	4134-14-099	4144	4144-14-099	4154	4154-14-099	4164	4164-14-099	4174	4174-14-099	4184	4184-14-099	4194	4194-14-099	4204	4204-14-099	4214	4214-14-099	4224	4224-14-099	4234	4234-14-099	4244	4244-14-099	4254	4254-14-099	4264	4264-14-099	4274	4274-14-099	4284	4284-14-099	4294	4294-14-099	4304	4304-14-099	4314	4314-14-099	4324	4324-14-099	4334	4334-14-099	4344	4344-14-099	4354	4354-14-099	4364	4364-14-099	4374	4374-14-099	4384	4384-14-099	4394	4394-14-099	4404	4404-14-099	4414	4414-14-099	4424	4424-14-099	4434	4434-14-099	4444	4444-14-099	4454	4454-14-099	4464	4464-14-099	4474	4474-14-099	4484	4484-14-099	4494	4494-14-099	4504	4504-14-099	4514	4514-14-099	4524	4524-14-099	4534	4534-14-099	4544	4544-14-099	4554	4554-14-099	4564	4564-14-099	4574	4574-14-099	4584	4584-14-099	4594	4594-14-099	4604	4604-14-099	4614	4614-14-099	4624	4624-14-099	4634	4634-14-099	4644	4644-14-099	4654	4654-14-099	4664	4664-14-099	4674	4674-14-099	4684	4684-14-099	4694	4694-14-099	4704	4704-14-099	4714	4714-14-099	4724	4724-14-099	4734	4734-14-099	4744	4744-14-099	4754	4754-14-099	4764	4764-14-099	4774	4774-14-099	4784	4784-14-099	4794	4794-14-099	4804	4804-14-099	4814	4814-14-099	4824	4824-14-099	4834	4834-14-099	4844	4844-14-099	4854	4854-14-099	4864	4864-14-099	4874	4874-14-099	4884	4884-14-099	4894	4894-14-099	4904	4904-14-099	4914	4914-14-099	4924	4924-14-099	4934	4934-14-099	4944	4944-14-099	4954	4954-14-099	4964	4964-14-099	4974	4974-14-099	4984	4984-14-099	4994	4994-14-099	5004	5004-14-099	5014	5014-14-099	5024	5024-14-099	5034</
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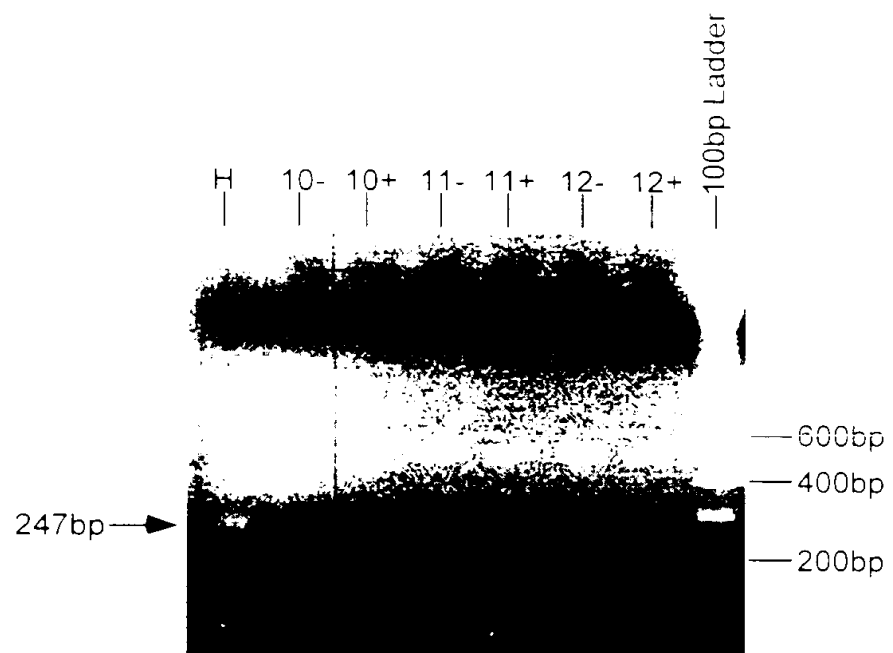


Figure 5. Expression of ADAMTS-E in cDNA from osteoarthritic cartilage.

[illegible]

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